## APPLICATION SERIAL NUMBER 10071648

## DOES NOT COMPLY WITH THE SEQUENCE RULES. See reasons below.

Paragraphs 0042 and 0044 disclose nucleotide sequences. No Sequence Listing has been provided.

Specification should be fully reviewed for occurrence of other sequences which meet the criteria for inclusion on a Sequence Listing.

SUPERVISORY PATENT EXAMINER
TECHNOLOGY CENTER 1800



# OIPE ROUTING SHEET APPLICATION TO BE DELIVERED TO: O I P E CP2-6C17 LOCA 0380

### Sequence Rule Compliance Review Item

X	CRF, paper copy of sequence listing, and statement that both are same missing
	CRF contains error(s) according to STIC Report
	CRF damaged or unreadable according to STIC Report
	CRF transferred from prior application is not compliant

Place an "X" in the appropriate box

DEBORAH J. REYNOLDS
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TECHNOLOGY CENTER 1600



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#### SEQUENCE LISTING

```
<110> DJAKIEW, DANIEL
 <120> METHODS FOR THE TREATMENT AND DIAGNOSIS OF PROSTATE
      CANCER BASED ON p75NTR TUMOR SUPRESSION
 <130> 082137/0280704
<140> 10/071,648
<141> 2002-02-11
<150> 60/268,940
<151> 2001-02-16
<160> 8
<170> PatentIn Ver. 2.1
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<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Primer
<400> 1
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<210> 2
<211> 20
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Primer
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                                                                    20
<210> 3
<211> 19
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Primer
<400> 3
agcttctcaa cggctctgc
                                                                    19
<210> 4
<211> 20
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<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Primer
<400> 4
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                                                                     20
<210> 5
<211> 21
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Primer
<400> 5
ccttctcccc acactgctag g
                                                                    21
<210> 6
<211> 20
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Primer
<400> 6
gcaagcatcc ccatctccac
                                                                    20
<210> 7
<211> 22
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Primer
<400> 7
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                                                                    22
<210> 8
<211> 23
<212> DNA
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<220>
<223> Description of Artificial Sequence: Primer
<400> 8
cagcagccag gatggagcaa tag
                                                                    23
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